

## SEQUENCE LISTING

- <110> Goulmy, Elsa
- <120> METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
- <130> 58994
- <140> 09/269,250
- <141> 1999-05-21
- <160> 38
- <170> PatentIn version 3.1
- <210> 1
- <211> 377
- <212> DNA
- <213> Human
- <400> 1
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- ggagggaggg acttggggag gctcagaagg\gagggaggct cagatggcag ggagggctgt
- gtggaagagg ccatgacagc taaggctctg aggatgtgt aggagtttgg tgggggagtc 180
- cctgagcgta cactggctca agagggtgcc cactttattt tttttaaagg atctgatggc 240
- aattaggagg gaaaggcaga ggaaatgtcc catgcacagg ctcagaaaca cggaaacaga
- gaatgcattt gggggccaag gtgtggggtg ccgctggtgt aggatgaagg catgacaacg

ccaggcagaa gggcaat

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2
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    20
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tggctctcac cgtcatgcag
    20
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      20
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      20
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cttaaggagt gtgtgctgca
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<211>
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gctgtcatgg cctcttccac
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       Artificial Sequence
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gtgtgctgca tgacggtg
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Val Leu Arg Asp Asp Leu Leu Glu Ala
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<222>
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gtg ctg cat gac gac ctc ctt gag gcc
Val Leu His Asp Asp Leu Leu Glu Ala
                5
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PRT

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                 5
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<211>
       23
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<220>
<223>
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gtgttgcgtg acggtgagag cca
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<210> 22
<211>
      37
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<223>
      Description of Artificial Sequence: Exon Fragments
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ctcactccga ctctccccag cagacctcct tgaggcc
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       23
<211>
      33
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Primer
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ccggcatgga cgtcgtcgag gacatctccc atc
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<211>
      30
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    30
<210> 25
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<400> 25
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Glu Cys Val Leu Arg Asp Asp Leu Leu Glu Ala Arg Arg
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Glu Cys Val Leu Arg Asp Asp Leu Leu Glu Ala Arg Arg
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       27
       39
<211>
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<223> Description of Artificial Sequence: PCR Product
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Glu Cys Val Leu His Asp Asp Leu Leu Glu Ala Arg Arg
                5
                                    10
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      28
<211> 13
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Glu Cys Val Leu His Asp Asp Leu Leu Glu Ala Arg Arg
<210>
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<211>
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       Human
<220>
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       Xaa represents a histidine (H) or an arginine (R) residue
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Val Leu Xaa Asp Asp Leu Leu Glu Ala
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<212> DNA
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gacgtcgtcg aggacatctc ccat
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gaaggccaca gcaatcgtct ccagg
    25
<210> 33
<211> 30
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<223> Description of Artificial Sequence: Primer
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       33
ccttgagaaa cttaaggagt gtgtgctgca
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ccttgagaaa cttaaggagt gtgtgttgcg 30

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Glu Cys Val Leu Arg Asp Asp Leu Leu Glu Ala Arg Arg Glu Cys Val

1

5

10

15

ctg cat gac gac ctc ctt gag gcc cgc cgc 78

Leu His Asp Asp Leu Leu Glu Ala Arg Arg

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25

<210> 36

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Glu Cys Val Leu Arg Asp Asp Leu Leu Glu Ala Arg Arg Glu Cys Val
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Leu His Asp Asp Leu Leu Glu Ala Arg Arg 20 25

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Tyr Xaa Thr Asp Arg Val Met Thr Val

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Val Leu His Asp Leu Leu Glu Ala 1 5